

References

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- NOTE: (Medline: 95314457) The anti-V3 Ab titre in patient serum was generally low against autologous virus isolated later than the serum sample, in contrast to a higher titre against peptides corresponding to virus isolated earlier than the serum sample. The authors conclude that the V3 domain is subject to immunoselection in vivo, and that V3 on early field virus is less accessible to NAb than the V3 loop on laboratory strains.
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- NOTE: (Medline: 94016886) Three closely related clones were derived from a neutralization resistant IIIB isolate that had been passaged in a chimpanzee. gp41 mutations were shown to profoundly alter the ability

of V3 loop MAbs 5023 and 178.1 to neutralize. Critical substitutions in gp41 were 668 and 675, close to the immunogenic domain 662-668, or ELDKWS. Less profound inhibition was observed for the anti-CD4 binding site MAb GP13.

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NOTE: (Medline: 96392164) A panel of anti-gp41 human Fab fragments were generated by panning phage display antibody libraries prepared from HIV-1 positive donors with rgp41. Fabs tended to be directed against three epitopes, designated clusters I-III. None were neutralizing. A common CDR3 motif was found in several of the heavy chain sequences.

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NOTE: (Medline: 96275629) Because N-linked glycans on viral glycoproteins can protect otherwise accessible neutralization epitopes of the viral envelope from neutralizing antibodies, the aim of this study was to explore the possibility of achieving a more broadly neutralizing immune response with a gp160 depleted of three N-linked glycans in the CD4-binding domain. Mutant and wild type gp160 were formulated into immunostimulating complexes (iscoms), and guinea pigs were vaccinated. Both preparations induced high serum antibody response to native gp120 and V3 peptides. The sera from animals immunized with the mutated glycoprotein lacking CD4 glycosylation sites did not neutralize nonrelated HIV strains better than did sera from animals immunized with wild type glycoprotein, but animals immunized with mutant gp160 neutralized mutant virus better than wild type virus, and vice versa.

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NOTE: (Medline: 95303114) Given the high degree of sequence variability of the V3 loop, the humoral response to this region tends to be type specific. An anti-idiotypic antibody could broaden the anti-V3 antibody polyclonal response in BALB/c mice relative to the original Ab used to generate the anti-idiotypic response. A synthetic peptide derived from the V3 determinant of HIV-1 MN induced an antibody response to multiple HIV-1 strains, but the extent of this cross-reactivity to be inversely correlated with the binding affinity to V3 MN peptide.

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NOTE: (Medline: 94322004) BALB/c mice were immunized with baculovirus expressed gp160 or gp120, and 15 MAbs were generated. No MAbs generated in this study neutralized reference strains, using a tetrazolium-based cytotoxicity assay to test for neutralization. Ten of the Mabs were mapped by peptide ELISA, and seven reacted with the C1 region, one with V2, one with V4, and one with the C-terminal end.

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NOTE: Medline: 94347460 A panel of 33 human monoclonal antibodies were produced. Linear epitopes for some of this set of MAbs were mapped using peptide ELISA. Linear epitopes were mapped in gp41, and a single epitope was mapped in p24. While multiple gp120 specific MAbs were generated, all seemed to be conformational or carbohydrate dependent, or both.

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NOTE: (Medline: 94340180) MAb F105 was administered intravenously to four cynomolgus monkeys. At 15 days post-dose, total serum F105 was 230 +/- 79 µg/ml and F105 was immunoreactive with cells infected with the MN and IIIB strains of HIV-1 as determined by flow cytometry.

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NOTE: (Medline: 96386562) The MAb 2F5 was infused into two chimpanzees which were then given an intravenous challenge with a primary HIV-1 isolate – both became infected, but with delayed detection and prolonged decrease in viral load relative to controls, indicating that preexisting, neutralizing antibodies (passively administered or actively elicited) affect the course of acute-phase virus replication and can be influential after the Ab can longer be detected in the peripheral circulation.

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NOTE: (Medline: 94211861) 2F5 is capable of neutralizing a broad range of primary isolates and lab strains. Susceptibility to neutralization was dependent on presence of a conserved antibody binding site. Kinetic studies were done, and 2F5 has a very long $t_{1/2}$ of dissociation, 156 minutes for gp41. The authors point out that LDKW core is present in highly diverged international isolates.

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NOTE: (Medline: 94240806) Antibodies against GalCer can block infection of CD4-negative cells from the brain and colon that are susceptible to HIV infection. This paper explores the ability of a panel to MAbs to inhibit binding of gp120 to GalCer, and also of the binding of GalCer to inhibit MAb-gp120 interaction. MAbs to the V3 loop and GalCer showed mutual inhibition of binding to gp120, and anti-CD4 binding site MAbs showed reduced inhibition. N- and C-terminal MAbs didn't influence GalCer binding.

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NOTE: (Medline: 96392163) Mice were injected with the gp120 in different configurations: free, complexed with CD4, and as an immunocomplex bound to a V3 loop MAb (M77) of the protein. Polyclonal sera, as well as of monoclonal antibodies produced in each case, were analyzed. The free gp120 and gp120-CD4 complex immunogens stimulated a responses were directed mainly toward conformational epitopes, but gp120 immunocomplexed with MAb M77 also produced numerous and varied MAbs directed toward linear epitopes that were presumably inaccessible on the gp120, gp120-CD4 proteins.

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NOTE: (Medline: 95121798) This paper describes the characterization of five antibodies that bind M77-epitopes that are only revealed upon M77-gp120 interaction.

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NOTE: (Medline: 95373192) To explore the immunogenicity of regions of gp120 that are exposed due to conformational changes in gp120 upon CD4 binding, CD4 was covalently linked to gp120 and this complex was used as an immunogen for BALB/c mice. Two MABs were produced, both of which bind preferentially to the gp120-CD4 complex, and are conformational. Competition assays indicate these MABs bind to epitopes that are recognized by sera from HIV-1 infected humans.

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NOTE: Medline: 94064668. The MAb M77 cannot neutralize a virus isolated from a IIIB infected lab-worker that has a single point mutation in the defined linear epitope. M77 cannot bind to the mutant native gp120, but can bind to a peptide that carries the substitution.

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NOTE: (Medline: 95114416) A panel of Fabs was obtained from a library prepared from the bone marrow of a long-term asymptomatic HIV-1 seropositive male donor. Four Fabs recognize the CD4BS. An additional four Fabs were retrieved after epitope masking gp120 with the CD4BS Fabs at the screening stage. 3/4 of these Fabs represent a V2 dependent conformational epitope.

- [Ditzel et al.(1997)] H. J. Ditzel, P. W. Parren, J. M. Binley, J. Sodroski, J. P. Moore, C. F. B. 3rd, & D. R. Burton. Mapping the protein surface of human immunodeficiency virus type 1 gp120 using human monoclonal antibodies from phage display libraries. *J Mol Biol* **267**:684–95, 1997.

NOTE: (Medline: 97272001), (Genbank: U82767 U82768 U82769 U82770 U82771 U82772 U82942 U82943 U82944 U82945 U82946 U82947 U82948 U82949 U82950 U82951 U82952 U82961 U82962) Recombinant monoclonal antibodies from phage display libraries provide a method for Env surface epitope mapping. Diverse epitopes are accessed by presenting gp120 to the library in different forms, such as sequential masking of epitopes with existing MABs or sCD4 prior to selection or by selection on peptides. Fabs identified by these methods have specificities associated with epitopes presented poorly on native multimeric envelope.

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NOTE: (Medline: 96014959) Eleven labs tested the 6 human MAbs 1125H, TH9, 4.8D, 257-D-IV, TH1, 2F5, and also HIVIG for neutralization of MN, JRCSF, the two B clade primary isolates 301657 and THA/92/026, and the D clade isolate UG/92/21. 2F5 was the most broadly neutralizing, better than HIVIG. The other MAbs showed limited neutralization of only MN (anti-CD4BS MAbs 1125H, TH9, and 4.8D), or MN and JRCSF (anti-V3 MAbs 257-D-IV and TH1).

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NOTE: (Medline: 97275172) Five laboratories evaluated neutralization of nine primary B clade isolates by a coded panel of seven human MAbs to HIV-1 subtype B envelope. IgG1b12, 2G12, 2F5 showed potent and broadly cross-reactive neutralizing ability; F105, 447/52-D, 729-D, 19b did not neutralize the primary isolates.

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NOTE: Medline: In a study of the repertoire of response to oligomeric versus monomeric Env protein, 138 murine MAbs were generated in response to an immunogen that was a gp120/bp41 oligomeric molecule that was not cleaved due to a mutation in the cleavage site. The oligomeric molecule was found to elicit a response that was very different than the monomer. Most MAbs were conformational, many were to gp41 or if in gp120, to the CD4 BS. Few MAbs to linear V3 epitopes were produced in response to oligomeric protein, though this was a common specificity in response to immunization with gp120 monomeric protein.

- [Eaton et al.(1994)] A. M. Eaton, K. E. Ugen, D. B. Weiner, T. Wildes, & J. A. Levy. An anti-gp41 human monoclonal antibody that enhances HIV-1 infection in the absence of complement. *AIDS Res Hum Retroviruses* **10**:13–18, 1994.

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NOTE: (Medline: 94107594) In this paper, three anti-HIV-1 gp41 specific MAbs were found to react with astrocytes: 98-6, 167-7 and 15G1. Reactive astrocytes in the hippocampus were most prominently involved, and the antibodies stained no other cell type in the brain, kidney or liver. All three mapped to a conformationally dependent epitope between aa 644-663.

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NOTE: (Medline: 97213943) To test whether antibody neutralization of HIV-1 primary isolates is correlated with the affinities for the oligomeric envelope glycoproteins, JRFL was used as a model primary virus and a panel of 13 human MAbs were evaluated for: half-maximal binding to rec monomeric JRFL gp120; half-maximal binding to oligomeric - JRFL Env expressed on the surface of transfected 293 cells; and neutralization of JRFL in a PBMC-based neutralization assay. Antibody affinity for oligomeric JRFL Env but not monomeric JRFL gp120 correlated with JRFL neutralization.

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NOTE: (Medline: 92114188) Two anti-envelope V2 antibodies were raised that neutralize virus in either a conformation dependent (G3-136) or conformation independent (BAT085) manner. G3-136 has diminished reactivity with deglycosylation or DTT reduced gp120, and sCD4 inhibits binding in a competition assay; BAT085 is not sensitive to these alterations in gp120.

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NOTE: (Medline: 96183910) Virus direct from plasma from six HIV-1 infected individuals was used for neutralization assay. MAb 19b could neutralize 2/6 plasma samples, while MAb IgG1b12 could neutralize 5/6 plasma samples. CD4-based molecules were also tested: CD4-IgG2 was effective in the *ex vivo* assay, but sCD4 was not. Thus, MAbs IgG1b12 and CD4-IgG2 have broad and potent *in vitro* and *ex vivo* neutralizing activities.

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NOTE: (Medline: 95271056) Passive protection against HIV-1 LAI with MAb BAT123 was achieved in SCID mice reconstituted with human peripheral blood lymphocytes (hu-PBL-SCID) BAT123 is specific for the V3 loop gp120 of HIV-1 LAI. Animals were protected against subsequent infection with LAI strain, but not other virus strains, when BAT123 was given 1 hour before virus inoculation, or up to 4 hours post-exposure. No therapeutic effect was observed when BAT123 was administered after infection had been established.

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NOTE: (Medline: 93387619) MAbs generated to a sCD4-gp120 complex, and the potential usefulness for vaccine design of epitopes specifically in the complex is discussed.

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NOTE: (Medline: 94188714) Crystal structure of V3 loop peptides bound to Fabs 59.1 and 50.1 was determined. The GPGRF motif forms a double turn.

- [Glaser & Hausdorf(1996)] R. W. Glaser & G. Hausdorf. Binding kinetics of an antibody against HIV p24 core protein measured with real-time biomolecular interaction analysis suggest a slow conformational change in antigen p24. *J Immunological Methods* **189**:1–14, 1996.

NOTE: (Medline: 96163539) The MAb CD-4/1 and p24 have unusual biphasic kinetics of association.

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NOTE: (Medline:).

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NOTE: (Medline: 93059712).

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NOTE: (Medline: 89160828) This paper described immortalization of B-cells from HIV-1 positive individuals with Epstein-Barr virus, to produce seven stable antibody producing cell lines.
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NOTE: (Medline: 95056063) Detailed characterization of the MAb 697-D.
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IgG1b12, including several non-B clade international isolates. Neutralization of a primary isolate with MAb IgG1b12 did not require continuous exposure to the antibody. A complete IgG1 molecule of a selected b12 FAb mutant with a 400-fold increase in affinity was assembled and evaluated in the infectivity reduction assay in comparative studies with the parent IgG1b12 antibody. The mutant did not retain the level of primary isolate neutralization potency of IgG1b12, despite the increase in affinity for gp120.

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NOTE: (Medline: 9168233) Multiple combinations of MAbs were tested for their ability to synergize neutralization of a SHIV construct containing HIV IIIB env. All of the MAb combinations tried were synergistic, suggesting such combinations may be useful for passive immunotherapy or immunoprophylaxis. Because SHIV can replicate in rhesus macaques, such approaches can potentially be studied in an *in vivo* monkey model.

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NOTE: (Medline: 93152284) The MAb M38 binds to the carboxy terminus of gp120, in a gp41 binding region. This MAb was used to create an anti-idiotypic MAb, 9G5A, which can bind to gp41 at the base of the cysteine loop. The binding domains of these two monoclonals are consistent with the C5 domain of gp120 being able to bind to the gp41 cysteine loop. The MAb M38 also binds to human HLA molecules, in antigenic homology or possibly molecular mimicry.

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NOTE:AIDSLINE: 1116291Abstract: VIIIth International Conference on AIDS, Florence, Italy, proceedings.

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NOTE: (Medline: 97456478) HIVIG derived from the plasma of HIV-1-infected donors, and MAbs 2F5 and 2G12 were tested against a panel of 15 clade B HIV-1 isolates, using a single concentration that is achievable in vivo (HIVIG, 2,500 microg/ml; MAbs, 25 microg/ml). While the three antibody reagents

neutralized many of the viruses tested, potency varied. The virus neutralization achieved by double or triple combinations was generally equal to or greater than that predicted by the effect of individual antibodies. and the triple combination was shown to be synergistic and to have the greatest breadth and potency. Passive immunotherapy for treatment or prophylaxis of HIV-1 should consider mixtures of these potent neutralizing antibody reagents.

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NOTE: (Medline: 93019072) Two MAbs are described that bind to a highly conserved region in p24, with antigenic conservation between FIV, SIV and HIV-1. The authors suggest this might be an immunodominant domain.

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NOTE: (Medline: 8661395) Chimeric viruses for HXB2 with primary isolate gp120 gave identical patterns of cell tropism and cytopathicity with the original primary viruses. Sera that were unable to neutralize the primary isolates were in some cases able to neutralize chimeric viruses, indicating that some of the neutralizing epitopes were in gp41.

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NOTE: (Medline: 93323237) Substitutions in the V2 loop can result in complete dissociation of gp120 and gp41, suggesting alterations in V2 can affect subunit assembly. Other substitutions allowed gp120-gp41 association and expression, but inhibited viral entry or syncytia. Binding of some neutralizing MAbs was altered by V2 substitutions. For MAb CRA-4, changes at residues 191/192/193 (YSL/GSS), and for 11/68b, changes at residues 183/184 (PI/SG), within V2, and for both MAbs a position 435 (Y/H) change in C4, abrogate binding. These MAbs can bind to V1 and V2 domains in the absence of C4 domain, so the C4 substitution probably results in conformational change.

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NOTE: (Medline: 97404700) A JRCSF resistant variant was selected by culturing in the presence of IgG1b12. The resistant virus remained sensitive to 2G12 and 2F5 and to CD4-IgG, encouraging for the possibility of combination therapy.

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NOTE: (Medline: 90274915).

[Moore et al.(1994a)] J. P. Moore, Y. Cao, D. D. Ho, & R. A. Koup. Development of the anti-gp120 antibody response during seroconversion to human immunodeficiency virus type 1. *J Virol* **68**:5142–5155, 1994a.

NOTE: (Medline: 94309181) Three seroconverting individuals were studied. The earliest detectable anti-gp120 antibodies were both conformational and anti-V3 loop, and could be detected only after the peak viremia has passed. No uniform pattern of autologous neutralizing anti-CD4BS or anti-V3 MAbs was observed.

[Moore et al.(1995a)] J. P. Moore, Y. Cao, L. Qing, Q. J. Sattentau, J. Pyati, R. Koduri, J. Robinson, C. F. Barbas III, D. R. Burton, & D. D. Ho. Primary isolates of human immunodeficiency virus type I are relatively resistant to neutralization by monoclonal antibodies to gp120, and their neutralization is not predicted by studies with monomeric gp120. *J Virol* **69**:101–109, 1995a.

NOTE: (Medline: 95074853) A panel of anti-gp120 MAbs and sera from HIV-1 infected individuals was tested for its ability to neutralize primary isolates. Most MAbs bound with high affinity to gp120 monomers from the various isolates, but were not effective at neutralizing. The MAb IgG1b12, which binds to a discontinuous anti-CD4 binding site epitope, was able to neutralize most of the primary isolates.

[Moore & Ho(1993)] J. P. Moore & D. D. Ho. Antibodies to discontinuous or conformationally sensitive epitopes on the gp120 glycoprotein of human immunodeficiency virus type 1 are highly prevalent in sera of infected humans. *J Virol* **67**:863–875, 1993.

NOTE: (Medline: 93124581) CD4BS antibodies are prevalent in HIV-1-positive sera, while neutralizing MAbs to C4, V2, and V3 and MAbs to linear epitopes are less common. Most linear epitope MAbs in human sera are directed against the V3 region, and cross-reactive MAbs tend to be directed against discontinuous epitopes.

[Moore & Ho(1995)] J. P. Moore & D. D. Ho. HIV-1 neutralization: the consequences of adaptation to growth on transformed T-cells. *AIDS* **9 suppl A**:S117–S136, 1995.

NOTE: (Medline: 96416784) This review considers the relative importance of a neutralizing antibody response for the development of a vaccine, and for disease progression during the chronic phase of HIV-1 infection. It suggests that T-cell immunity may be more important. The distinction between MAbs that can neutralize primary isolates, and those that are effective at neutralizing only laboratory adapted strains is discussed in detail. Alternative conformations of envelope and non-contiguous interacting domains in gp120 are discussed. The suggestion that soluble monomeric gp120 may serve as a viral decoy that diverts the humoral immune response *in vivo* is put forth.

[Moore et al.(1994b)] J. P. Moore, F. E. McCutchan, S.-W. Poon, J. Mascola, J. Liu, Y. Cao, & D. D. Ho. Exploration of antigenic variation in gp120 from clades A through F of human immunodeficiency virus type 1 by using monoclonal antibodies. *J Virol* **68**:8350–8364, 1994b.

NOTE: (Medline: 95056067) Four of five anti-V3 MAbs were slightly cross-reactive within clade B, but not very reactive outside clade B. Two discontinuous CD4 binding site Mabs appear to be pan-reactive. Anti-V2 MAbs were only sporadically reactive inside and outside of clade B.

[Moore et al.(1990)] J. P. Moore, J. A. McKeating, R. A. Weiss, & Q. J. Sattentau. Dissociation of gp120 from HIV-1 virions induced by soluble CD4. *Science* **250**:1139–1142, 1990.

NOTE: AIDSLINE: 91068008.

[Moore et al.(1994c)] J. P. Moore, Q. J. Sattentau, R. Wyatt, & J. Sodroski. Probing the structure of the human immunodeficiency virus surface glycoprotein gp120 with a panel of monoclonal antibodies. *J Virol* **68**:469–484, 1994c.

NOTE: (Medline: 94076440). This study compared a large number of MAbs that bind to linear epitopes of gp120, and compared binding affinities for: i) native and SDS-DDT denatured gp120, (clone BH10 of the LAI isolate expressed in CHO cells); ii) recombinant gp120 lacking the V1, V2, V3 loops; iii) a panel of 20 mer peptides; iv) a panel of gp120 mutants; and v) oligomeric versus monomeric gp120. The binding ratio of native versus denatured monomeric gp120 is included in the table in this database. These numbers should be considered with the following points in mind: a continuous epitope may be partially exposed on the surface; and a preparation of rgp120 is not homogeneous and contains fully folded, partly denatured, and some completely unfolded species, so the conformation of what is considered to be a native protein will not only reflect fully folded gp120. The authors suggest that a fivefold increase in the affinity for a MAb binding to denatured versus native gp120 indicates that the epitope is inaccessible in the native form. We also have included here information extracted from Moore et al's list of the gp120 mutations that reduced the binding of a particular MAb. In mapping of exposed regions of gp120, C2, C3, and C5 domain epitopes were found to bind preferentially to denatured gp120. V1, V2 and V3, part of C4, and the extreme carboxy terminus of C5 were exposed on the native monomer. In the oligomeric form of the molecule, only V2, V3 and part of C4 are well exposed as continuous epitopes.

[Moore et al.(1993a)] J. P. Moore, Q. J. Sattentau, H. Yoshiyama, M. Thali, M. Charles, N. Sullivan, S.-W. Poon, M. S. Fung, F. Traincard, M. Pinkus, G. Robey, J. E. Robinson, D. D. Ho, & J. Sodroski. Probing the structure of the V2 domain of human immunodeficiency virus type 1 surface glycoprotein gp120 with a panel of eight monoclonal antibodies: human immune response to the V1 and V2 domains. *J Virol* **67**:6136–6151, 1993a.

NOTE: (Medline: 93381817).

[Moore & Sodroski(1996)] J. P. Moore & J. Sodroski. Antibody cross-competition analysis of the human immunodeficiency virus type 1 gp120 exterior envelope glycoprotein. *J Virol* **70**:1863–1872, 1996.

NOTE: AIDSLINE: 96190589 46 anti-gp120 monomer MAbs were used to create a competition matrix, and MAb competition groups were defined. The data suggests that there are two faces of the gp120 glycoprotein: a face occupied by the CD4BS, which is presumably also exposed on the oligomeric envelope glycoprotein complex, and a second face which is presumably inaccessible on the oligomer and interacts with a number of nonneutralizing antibodies.

[Moore et al.(1993b)] J. P. Moore, M. Thali, B. A. Jameson, F. Vignaux, G. K. Lewis, S.-W. Poon, M. S. Fung, P. J. Durda, L. Akerblom, B. Wahren, D. D. Ho, Q. J. Sattentau, & J. Sodroski. Immunochemical analysis of the gp120 surface glycoprotein of human immunodeficiency virus type 1: Probing the structure of the C4 and V4 domains and the interaction of the C4 domain with the V3 loop. *J Virol* **73**:4785–4796, 1993b.

NOTE: Medline: 93323221. General observations: C4 and V3 MAbs are sensitive to the way the epitopes are presented, and this sensitivity cannot be correlated to peptide binding. Some V3-C4 domain interaction was indicated based on mutation and interference studies.

[Moore et al.(1995b)] J. P. Moore, A. Trkola, B. Korber, L. J. Boots, J. A. Kessler II, F. E. McCutchan, J. Mascola, D. D. Ho, J. Robinson, & A. J. Conley. A human monoclonal antibody to a complex epitope

in the V3 region of gp120 of human immunodeficiency virus type 1 has broad reactivity within and outside clade B. *J Virol* **69**:122–130, 1995b.

NOTE: (Medline: 95074855) The epitope was defined as including amino acids on both sides of the loop of the V3 loop: -I—G—FY-T, where the G is the second G of the GPGR tip of the loop. This antibody bound well to gp120 molecules from clades A,B,C,E, and F, when the critical amino acids were present. Binding did not parallel neutralization however; 19b could produce a 50-fold reduction of infectivity in some primary B isolates, and in C clade isolates at low virus input concentrations, but not in isolates from all clades where binding could occur (A,E, and F).

[Moore et al.(1994d)] J. P. Moore, R. L. Willey, G. K. Lewis, J. Robinson, & J. Sodroski. Immunological evidence for interactions between the first, second and fifth conserved domains of the gp120 surface glycoprotein of human immunodeficiency virus type 1. *J Virol* **68**:6836–6847, 1994d.

NOTE: (Medline: 95018590) Mutation 267N/Q in C2 region results in exposing the carboxy-terminal end gp120.

[Moore et al.(1993c)] J. P. Moore, H. Yoshiyama, D. D. Ho, J. E. Robinson, & J. Sodroski. Antigenic variation in gp120s from molecular clones of HIV-1 LAI. *AIDS Res Hum Retroviruses* **9**:1185–1193, 1993c.

NOTE: AIDSLINE: 94190623 The binding MAbs to four molecular clones of HIV-1 LAI: HxB2, HxB3, Hx10, and NL4-3, was measured. Despite the close relationship between these clones, there is considerable variation in their antigenic structure, judged by MAb reactivities to the V2, V3, and C4 domains and to discontinuous epitopes. Small variations in sequence can profoundly affect recognition of gp120 by all five groups of defined anti-gp120 neutralizing antibodies.

[Moran et al.(1993)] M. J. Moran, J. S. Andris, Y. Matsumoto, J. D. Capra, & E. M. Hersh. Variable region genes of anti-HIV human monoclonal antibodies: Non-restricted use of the V gene repertoire and extensive somatic mutation. *Mol Immunol* **30**:1543–1551, 1993.

NOTE: (Medline: 94049845) Sequenced variable regions from four human anti-HIV-1 MAbs: anti-gp120 13, S1-1 and HBW4; and anti-gp41 No.86. Extensive somatic mutation was observed and under-representation of V_H III usage.

[Muller et al.(1991)] S. Muller, H.-T. Wang, S.-V. Kaveri, S. Chattopadhyay, & H. Kohler. Generation and specificity of monoclonal anti-idiotypic antibodies against human HIV-specific antibodies. *J Immunol* **147**:933–941, 1991.

NOTE: (Medline: 91318181).

[Muster et al.(1995)] T. Muster, B. Ferko, A. Klima, M. Purtscher, A. Trkola, P. Schulz, A. Grassauer, O. G. Englehard, A. Garcia-Sastre, P. Palese, & H. Katinger. Mucosal model of immunization against human immunodeficiency virus type 1 with a chimeric influenza virus. *J Virol* **69**:6678–6686, 1995.

NOTE: (Medline: 96013760).

[Muster et al.(1994)] T. Muster, R. Guinea, A. Trkola, M. Purtscher, A. Klima, F. Steindl, P. Palese, & H. Katinger. Cross-neutralization activity against divergent human immunodeficiency virus type 1 isolates induced by the gp41 sequence ELDKWAS. *J Virol* **68**:4031–4034, 1994.

NOTE: (Medline: 94246751).

[Muster et al.(1993)] T. Muster, F. Steindl, M. Purtscher, A. Trkola, A. Klima, G. Himmler, F. Ruker, & H. Katinger. A conserved neutralizing epitope on gp41 of human immunodeficiency virus type 1. *J Virol* **67**:6642–6647, 1993.

NOTE: (Medline: 94016848) Peptides containing the amino acid sequence LDKWAS or DKWASL showed reduced reactivity. The peptides LELDKW and KWASLW showed no significant reaction. These data suggest that the epitope of the MAb 2F5 comprises the amino acid sequence ELDKWA, with DKWA being the core sequence.

[Myers et al.(1993)] R. Myers, T. Meiller, W. Falkler Jr., J. Patel, & J. Joseph. A human monoclonal antibody to a cryptic gp41 epitope on HIV-1 infected cells. *Abstr Gen Meet Am Soc Microbiol* **93**:444, 1993.

NOTE: Aidsline: 93291838 Abstract T70.

[Nakamura et al.(1992)] G. R. Nakamura, R. Byrn, K. Rosenthal, J. P. Porter, M. R. Hobbs, L. Riddle, D. J. Eastman, D. Dowbenko, T. Gregory, B. M. Fendly, & P. W. Berman. Monoclonal antibodies to the extracellular domain of HIV-1 IIIB gp160 that neutralize infectivity, block binding to CD4, and react with diverse isolates. *AIDS Res Hum Retroviruses* **8**:1875–1885, 1992.

NOTE: (Medline: 93143997).

[Nakamura et al.(1993)] G. R. Nakamura, R. Byrn, D. M. Wilkes, J. A. Fox, M. R. Hobbs, R. Hastings, H. C. Wessling, M. A. Norcross, B. M. Fendly, & P. W. Berman. Strain specificity and binding affinity requirements of neutralizing monoclonal antibodies to the C4 domain of gp120 from human immunodeficiency virus type 1. *J Virol* **67**:6179–6191, 1993.

NOTE: (Medline: 93381821) Multiple CD4 binding domain antibodies are described; only one has a linear peptide reactivity (13H8). A V3 loop binding antibody is also described (1026).

[Nara et al.(1990)] P. L. Nara, L. Smit, N. Dunlop, W. Hatch, M. Merges, D. Waters, J. Kelliher, R. C. Gallo, P. J. Fischinger, & J. Goudsmit. Emergence of viruses resistant to neutralization by V3-specific antibodies in experimental human immunodeficiency virus type 1 IIIB infection of chimpanzees. *J Virol* **64**:3779–3791, 1990.

NOTE: (Medline: 90317876).

[Neurath & Strick(1990)] A. R. Neurath & N. Strick. Confronting the hypervariability of an immunodominant epitope eliciting virus neutralizing antibodies from the envelope glycoprotein of the human immunodeficiency virus type 1. *Mol Immunol* **27**:539–549, 1990.

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NOTE: (Medline: 97275172) A series of HIV-1 envelope glycoproteins from related primary virus isolates of different SI phenotypes, together with chimeras of these proteins, were tested in an envelope trans-complementation assay for their sensitivity to either antibody mediated inhibition or enhancement of HIV-1 entry. In contrast to the inhibition of HIV-1 entry, antibody mediated enhancement was not temperature dependent and could not be mediated by F(ab) fragments, implicating cross-linking as an important step. Enhancement or inhibition seemed to be determined by virus isolate rather than by the specificity of the antiserum used. 2F5 was the only MAb that inhibited the entry of all viruses.

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NOTE: (Medline: 91333022) Six MAbs with linear epitopes were mapped. These Abs could only bind to HIV-infected cells that had been permeabilized with acetone. Only G11g1 and G11h3, two antibodies that did not bind to peptides, but only to intact p17, could react with live HIV-1 infected cells. These antibodies were not neutralizing.

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NOTE: Medline: 95074868. Anti-V2 linear and conformation dependent MAbs were studied. All V2 Abs studied could bind IIIB, but failed to neutralize non-clonal stocks. Epitope exposure is different in rgp120 compared to native gp120. HXB2 V2-MAb neutralization escape mutants were sequenced.

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NOTE: AIDSLINE: 94133079 Pseudotypes were formed with HIV and HTLV-I. MAb 9284, directed at the V3 loop of gp120, failed to inhibit the infection of CD-4 negative cells with pseudotypes, but anti-HTLV serum did inhibit infection. HIV and HTLV-I appear to induce common carbohydrate neutralizing epitopes.

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NOTE: (Medline: 93100837) This study looked at the ability of 16 human MAbs to activate complement. MAbs directed against the V3 region could induce C3 deposition on infected cells and virolysis of free virus, but antibodies to the CD4BS and C-terminal region and two regions in gp41 could induce no complement mediated effects. Pre-treatment with sCD4 could increased complement-mediated effects of anti-gp41 MAbs, but decreased the complement-mediated effects of V3 MAbs. Anti-gp41 MAbs were able to affect IIIB but not MN virolysis, suggesting spontaneous shedding of gp120 on IIIB virions exposes gp41 epitopes. IgG isotype did not appear to have an effect on virolysis or C3 deposition.

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NOTE: (Medline: 96136775) HIV and HIV-infected cells are not subject to efficient complement-mediated lysis, even in the presence of HIV-specific antibodies. HIV is intrinsically resistant to human complement. Decay accelerating factor (DAF) and human complement factor H (CFH), a humoral negative regulator of complement which binds to gp41 are critical for this resistance. MAb 2F5 can inhibit CHF binding and facilitate complement mediated lysis.

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NOTE: AIDSLINE: 95287498 Three gp120 molecules derived from primary isolates were compared to T-cell adapted lines HXBc2 and MN. Complementation experiments showed viral entry into peripheral blood mononuclear cell targets was five-fold less efficient for primary isolates. Anti-CD4 binding site neutralizing MAbs were far less potent against primary isolates, and the single anti-V3 MAb tested was 3-fold less potent. The differences in neutralization efficiency could not be attributed to differences in affinity for monomeric gp120, but were related to binding to the oligomeric complex. Enhanced infectivity of primary isolates was observed using sCD4 and MAb F105, which can neutralize T-cell adapted strains.

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NOTE: (Medline: 93267832) Recognition of neutralizing MAb G3-4 was altered by substitutions in 176 to 184 in the V2 loop. Some changes in the V2 loop can affect subunit assembly; other changes allow expression and CD4 binding but inhibit syncytium formation and viral entry, suggesting that V1/V2 may be involved in post receptor binding events.

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epitopes in gp120. N70-1.5 is a potent neutralizing MAb with no enhancing activity, while N70-2.3a doesn't neutralize and mediates enhancement of HIV-1 infection.

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NOTE: (Medline: 94118414) A T/A amino acid substitution at position 582 of gp41 conferred resistance to neutralization to 30% of HIV positive sera (Wilson et al. *J Virol* 64:3240-48 (1990)). Monoclonal antibodies that bound to the CD4 binding site were unable to neutralize this virus, but the mutation did not reduce the neutralizing capacity of a V2 region MAb G3-4, V3 region MABs, or gp41 neutralizing MAb 2F5.

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